



IFWO

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/749,104

DATE: 08/31/2004
TIME: 15:46:39

Input Set : A:\PTO.LM.txt
Output Set: N:\CRF4\08312004\J749104.raw

5 <110> APPLICANT: Shen Zhen Tsinghua Yuanxing Bio-Pharm Science & Technology Co., Ltd.
9 <120> TITLE OF INVENTION: Anti-Tumor Molecular Vaccine and Method of making thereof
13 <130> FILE REFERENCE: CGCNS31755
C--> 17 <140> CURRENT APPLICATION NUMBER: US/10/749,104
C--> 17 <141> CURRENT FILING DATE: 2003-12-29
17 <150> PRIOR APPLICATION NUMBER: CN02159602.6
18 <151> PRIOR FILING DATE: 2002-12-27
20 <160> NUMBER OF SEQ ID NOS: 27
24 <170> SOFTWARE: PatentIn version 3.1
28 <210> SEQ ID NO: 1
30 <211> LENGTH: 3633
32 <212> TYPE: DNA
34 <213> ORGANISM: EE(homo sapiens)
38 <220> FEATURE:
40 <221> NAME/KEY: CDS
42 <222> LOCATION: (1)..(3630)
44 <223> OTHER INFORMATION:

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50 Met Arg Pro Ser Gly Thr Ala Gly Ala Ala Leu Leu Ala Leu Leu Ala
51 1 5 10 15
53 gcg ctc tgc ccg gcg agt cggt gct ctg gag gaa aag aaa gtt tgc caa 96
54 Ala Leu Cys Pro Ala Ser Arg Ala Leu Glu Glu Lys Lys Val Cys Gln
55 20 25 30
57 ggc acg agt aac aag ctc acg cag ttg ggc act ttt gaa gat cat ttt 144
58 Gly Thr Ser Asn Lys Leu Thr Gln Leu Gly Thr Phe Glu Asp His Phe
59 35 40 45
61 ctc agc ctc cag agg atg ttc aat aac tgt gag gtg gtc ctt ggg aat 192
62 Leu Ser Leu Gln Arg Met Phe Asn Asn Cys Glu Val Val Leu Gly Asn
63 50 55 60
65 ttg gaa att acc tat gtg cag agg aat tat gat ctt tcc ttc tta aag 240
66 Leu Glu Ile Thr Tyr Val Gln Arg Asn Tyr Asp Leu Ser Phe Leu Lys
67 65 70 75 80
69 acc atc cag gag gtg gct ggt tat gtc ctc att gcc ctc aac aca gtg 288
70 Thr Ile Gln Glu Val Ala Gly Tyr Val Leu Ile Ala Leu Asn Thr Val
71 85 90 95
73 gag cga att cct ttg gaa aac ctg cag atc atc aga gga aat atg tac 336
74 Glu Arg Ile Pro Leu Glu Asn Leu Gln Ile Ile Arg Gly Asn Met Tyr
75 100 105 110
77 tac gaa aat tcc tat gcc tta gca gtc tta tct aac tat gat gca aat 384
78 Tyr Glu Asn Ser Tyr Ala Leu Ala Val Leu Ser Asn Tyr Asp Ala Asn
79 115 120 125
81 aaa acc gga ctg aag gag ctg ccc atg aga aat tta cag gaa atc ctg 432

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82 Lys Thr Gly Leu Lys Glu Leu Pro Met Arg Asn Leu Gln Glu Ile Leu
83      130      135      140
85 cat ggc gcc gtg cgg ttc agc aac aac cct gcc ctg tgc aac gtg gag      480
86 His Gly Ala Val Arg Phe Ser Asn Asn Pro Ala Leu Cys Asn Val Glu
87 145      150      155      160
89 agc atc cag tgg cgg gac ata gtc agc agt gac ttt ctc agc aac atg      528
90 Ser Ile Gln Trp Arg Asp Ile Val Ser Ser Asp Phe Leu Ser Asn Met
91      165      170      175
93 tcg atg gac ttc cag aac cac ctg ggc agc tgc caa aag tgt gat cca      576
94 Ser Met Asp Phe Gln Asn His Leu Gly Ser Cys Gln Lys Cys Asp Pro
95      180      185      190
97 agc tgt ccc aat ggg agc tgc tgg ggt gca gga gag gag aac tgc cag      624
98 Ser Cys Pro Asn Gly Ser Cys Trp Gly Ala Gly Glu Glu Asn Cys Gln
99      195      200      205
101 aaa ctg acc aaa atc atc tgt gcc cag cag tgc tcc ggg cgc tgc cgt      672
102 Lys Leu Thr Lys Ile Ile Cys Ala Gln Gln Cys Ser Gly Arg Cys Arg
103 210      215      220
105 ggc aag tcc ccc agt gac tgc tgc cac aac cag tgt gct gca ggc tgc      720
106 Gly Lys Ser Pro Ser Asp Cys Cys His Asn Gln Cys Ala Ala Gly Cys
107 225      230      235      240
109 aca ggc ccc cgg gag agc gac tgc ctg gtc tgc cgc aaa ttc cga gac      768
110 Thr Gly Pro Arg Glu Ser Asp Cys Leu Val Cys Arg Lys Phe Arg Asp
111      245      250      255
113 gaa gcc acg tgc aag gac acc tgc ccc cca ctc atg ctc tac aac ccc      816
114 Glu Ala Thr Cys Lys Asp Thr Cys Pro Pro Leu Met Leu Tyr Asn Pro
115 260      265      270
117 acc acg tac cag atg gat gtg aac ccc gag ggc aaa tac agc ttt ggt      864
118 Thr Thr Tyr Gln Met Asp Val Asn Pro Glu Gly Lys Tyr Ser Phe Gly
119 275      280      285
121 gcc acc tgc gtg aag aag tgt ccc cgt aat tat gtg gtg aca gat cac      912
122 Ala Thr Cys Val Lys Lys Cys Pro Arg Asn Tyr Val Val Thr Asp His
123 290      295      300
125 ggc tgc tgc gtc cga gcc tgt ggg gcc gac agc tat gag atg gag gaa      960
126 Gly Ser Cys Val Arg Ala Cys Gly Ala Asp Ser Tyr Glu Met Glu Glu
127 305      310      315      320
129 gac ggc gtc cgc aag tgt aag aag tgc gaa ggg cct tgc cgc aaa gtg      1008
130 Asp Gly Val Arg Lys Cys Lys Cys Glu Gly Pro Cys Arg Lys Val
131      325      330      335
133 tgt aac gga ata ggt att ggt gaa ttt aaa gac tca ctc tcc ata aat      1056
134 Cys Asn Gly Ile Gly Ile Gly Glu Phe Lys Asp Ser Leu Ser Ile Asn
135      340      345      350
137 gct acg aat att aaa cac ttc aaa aac tgc acc tcc atc agt ggc gat      1104
138 Ala Thr Asn Ile Lys His Phe Lys Asn Cys Thr Ser Ile Ser Gly Asp
139      355      360      365
141 ctc cac atc ctg cgg gtg gca ttt agg ggt gac tcc ttc aca cat act      1152
142 Leu His Ile Leu Pro Val Ala Phe Arg Gly Asp Ser Phe Thr His Thr
143      370      375      380
145 cct cct ctg gat cca cag gaa ctg gat att ctg aaa acc gta aag gaa      1200
146 Pro Pro Leu Asp Pro Gln Glu Leu Asp Ile Leu Lys Thr Val Lys Glu
  
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147	385	390	395	400	
149	atc aca ggg ttt ttg ctg att cag gct tgg cct gaa aac agg acg gac				1248
150	Ile Thr Gly Phe Leu Leu Ile Gln Ala Trp Pro Glu Asn Arg Thr Asp				
151	405	410	415		
153	ctc cat gcc ttt gag aac cta gaa atc ata cgc ggc agg acc aag caa				1296
154	Leu His Ala Phe Glu Asn Leu Glu Ile Ile Arg Gly Arg Thr Lys Gln				
155	420	425	430		
157	cat ggt cag ttt tct ctt gca gtc agc ctg aac ata aca tcc ttg				1344
158	His Gly Gln Phe Ser Leu Ala Val Val Ser Leu Asn Ile Thr Ser Leu				
159	435	440	445		
161	gga tta cgc tcc ctc aag gag ata agt gat gga gat gtt ata att tca				1392
162	Gly Leu Arg Ser Leu Lys Glu Ile Ser Asp Gly Asp Val Ile Ile Ser				
163	450	455	460		
165	gga aac aaa aat ttg tgc tat gca aat aca ata aac tgg aaa aaa ctg				1440
166	Gly Asn Lys Asn Leu Cys Tyr Ala Asn Thr Ile Asn Trp Lys Lys Leu				
167	465	470	475	480	
169	ttt ggg acc tcc ggt cag aaa acc aaa att ata agc aac aga ggt gaa				1488
170	Phe Gly Thr Ser Gly Gln Lys Thr Lys Ile Ile Ser Asn Arg Gly Glu				
171	485	490	495		
173	aac agc tgc aag gcc aca ggc cag gtc tgc cat gcc ttg tgc tcc ccc				1536
174	Asn Ser Cys Lys Ala Thr Gly Gln Val Cys His Ala Leu Cys Ser Pro				
175	500	505	510		
177	gag ggc tgc tgg ggc ccg gag ccc agg gac tgc gtc tct tgc cgg aat				1584
178	Glu Gly Cys Trp Gly Pro Glu Pro Arg Asp Cys Val Ser Cys Arg Asn				
179	515	520	525		
181	gtc agc cga ggc agg gaa tgc gtg gac aag tgc aac ctt ctg gag ggt				1632
182	Val Ser Arg Gly Arg Glu Cys Val Asp Lys Cys Asn Leu Leu Glu Gly				
183	530	535	540		
185	gag cca agg gag ttt gtg gag aac tct gag tgc ata cag tgc cac cca				1680
186	Glu Pro Arg Glu Phe Val Glu Asn Ser Glu Cys Ile Gln Cys His Pro				
187	545	550	555	560	
189	gag tgc ctg cct cag gcc atg aac atc acc tgc aca gga cgg gga cca				1728
190	Glu Cys Leu Pro Gln Ala Met Asn Ile Thr Cys Thr Gly Arg Gly Pro				
191	565	570	575		
193	gac aac tgt atc cag tgt gcc cac tac att gac ggc ccc cac tgc gtc				1776
194	Asp Asn Cys Ile Gln Cys Ala His Tyr Ile Asp Gly Pro His Cys Val				
195	580	585	590		
197	aag acc tgc ccg gca gga gtc atg gga gaa aac aac acc ctg gtc tgg				1824
198	Lys Thr Cys Pro Ala Gly Val Met Gly Glu Asn Asn Thr Leu Val Trp				
199	595	600	605		
201	aag tac gca gac gcc ggc cat gtg tgc cac ctg tgc cat cca aac tgc				1872
202	Lys Tyr Ala Asp Ala Gly His Val Cys His Leu Cys His Pro Asn Cys				
203	610	615	620		
205	acc tac gga tgc act ggg cca ggt ctt gaa ggc tgt cca acg aat ggg				1920
206	Thr Tyr Gly Cys Thr Gly Pro Gly Leu Glu Gly Cys Pro Thr Asn Gly				
207	625	630	635	640	
209	cct aag atc ccg tcc atc gcc act ggg atg gtg ggg gcc ctc ctc ttg				1968
210	Pro Lys Ile Pro Ser Ile Ala Thr Gly Met Val Gly Ala Leu Leu Leu				
211	645	650	655		

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213	ctg	ctg	gtg	gtg	gcc	ctg	ggg	atc	ggc	ctc	ttc	atg	cga	agg	cgc	cac	2016		
214	Leu	Leu	Val	Val	Ala	Leu	Gly	Ile	Gly	Leu	Phe	Met	Arg	Arg	Arg	His			
215								660			665			670					
217	atc	gtt	cg	gg	aag	cg	cc	ac	gt	cg	gg	at	cg	ga	gg	ct	2064		
218	Ile	Val	Arg	Lys	Arg	Thr	Leu	Arg	Arg	Leu	Leu	Gln	Glu	Arg	Glu	Leu			
219								675			680			685					
221	gtg	gag	cct	ctt	aca	ccc	agt	gga	gaa	gct	ccc	aa	caa	gct	ctc	tt	2112		
222	Val	Glu	Pro	Leu	Thr	Pro	Ser	Gly	Glu	Ala	Pro	Asn	Gln	Ala	Leu	Leu			
223								690			695			700					
225	agg	atc	tt	g	aag	gaa	act	gaa	ttc	aaa	aag	atc	aaa	gtg	ctg	ggc	2160		
226	Arg	Ile	Leu	Lys	Glu	Thr	Glu	Phe	Lys	Lys	Ile	Lys	Val	Leu	Gly	Ser			
227								705			710			715		720			
229	ggt	g	cg	tt	gg	ac	g	tg	tat	aag	g	ct	tgg	atc	cca	gaa	ggt	gag	2208
230	Gly	Ala	Phe	Gly	Thr	Val	Tyr	Lys	Gly	Leu	Trp	Ile	Pro	Glu	Gly	Glu			
231								725			730			735					
233	aaa	gtt	aaa	att	ccc	gtc	gct	atc	aag	gaa	tta	aga	gaa	gca	aca	tct	2256		
234	Lys	Val	Lys	Ile	Pro	Val	Ala	Ile	Lys	Glu	Leu	Arg	Glu	Ala	Thr	Ser			
235								740			745			750					
237	ccg	aaa	gg	cc	aa	g	aa	atc	ct	gat	gaa	gg	tac	gt	atg	gg	2304		
238	Pro	Lys	Ala	Asn	Lys	Glu	Ile	Leu	Asp	Glu	Ala	Tyr	Val	Met	Ala	Ser			
239								755			760			765					
241	gtg	gac	aa	cc	cc	cac	gt	tg	cg	ct	gg	atc	tg	ct	acc	tt	2352		
242	Val	Asp	Asn	Pro	His	Val	Cys	Arg	Leu	Leu	Gly	Ile	Cys	Leu	Thr	Ser			
243								770			775			780					
245	acc	gt	ca	ct	atc	ac	ca	gt	ct	at	cc	tt	gg	cc	tg	ct	2400		
246	Thr	Val	Gln	Leu	Ile	Thr	Gln	Leu	Met	Pro	Phe	Gly	Cys	Leu	Leu	Asp			
247								785			790			795		800			
249	tat	gt	cc	gg	aa	ca	aa	gac	aa	tat	gg	tc	ca	tg	ct	aa	2448		
250	Tyr	Val	Arg	Glu	His	Lys	Asp	Asn	Ile	Gly	Ser	Gln	Tyr	Leu	Leu	Asn			
251								805			810			815					
253	tgg	tgt	gt	ca	atc	gca	aa	gg	at	aa	tc	tt	ga	ga	cgt	cg	2496		
254	Trp	Cys	Val	Gln	Ile	Ala	Lys	Gly	Met	Asn	Tyr	Leu	Glu	Asp	Arg	Arg			
255								820			825			830					
257	ttg	tg	tg	ca	cg	gac	ct	g	cc	gg	aa	ac	ta	ct	gt	aa	aca	cc	2544
258	Leu	Val	His	Arg	Asp	Leu	Ala	Ala	Arg	Asn	Val	Leu	Val	Lys	Thr	Pro			
259								835			840			845					
261	cag	ca	gt	aa	atc	ac	aa	gat	ttt	gg	ct	gg	aa	aa	ct	ct	gg	2592	
262	Gln	His	Val	Lys	Ile	Thr	Asp	Phe	Gly	Leu	Ala	Lys	Leu	Leu	Gly	Ala			
263								850			855			860					
265	gaa	gag	aa	aa	ta	ca	ca	aa	gt	cc	tt	gg	aa	aa	ct	ct	gg	2640	
266	Glu	Glu	Lys	Glu	Tyr	His	Ala	Glu	Gly	Gly	Lys	Val	Pro	Ile	Lys	Trp			
267								865			870			875		880			
269	at	g	ca	tt	g	aa	tc	at	ca	aa	tc	ca	ca	ag	gt	at	2688		
270	Met	Ala	Leu	Glu	Ser	Ile	Leu	His	Arg	Ile	Tyr	Thr	His	Gln	Ser	Asp			
271								885			890			895					
273	gtc	tgg	agc	ta	gg	gg	gt	gg	ac	tt	gg	at	gg	ac	tt	gg	2736		
274	Val	Trp	Ser	Tyr	Gly	Val	Thr	Val	Trp	Glu	Leu	Met	Thr	Phe	Gly	Ser			
275								900			905			910					
277	aag	cc	ta	g	ac	atc	c	cc	gg	atc	tcc	tcc	atc	ct	gg	gg	2784		

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282	Lys	Gly	Glu	Arg	Leu	Pro	Gln	Pro	Pro	Ile	Cys	Thr	Ile	Asp	Val	Tyr	
283		930			935					940							
285	atg	atc	atg	gtc	aag	tgc	tgg	atg	ata	gac	gca	gat	agt	cgc	cca	aag	2880
286	Met	Ile	Met	Val	Lys	Cys	Trp	Met	Ile	Asp	Ala	Asp	Ser	Arg	Pro	Lys	
287	945		950			955			960								
289	ttc	cgt	gag	ttg	atc	atc	gaa	ttc	tcc	aaa	atg	gcc	cga	gac	ccc	cag	2928
290	Phe	Arg	Glu	Leu	Ile	Ile	Glu	Phe	Ser	Lys	Met	Ala	Arg	Asp	Pro	Gln	
291		965			970					975							
293	cgc	tac	ctt	gtc	att	cag	ggg	gat	gaa	aga	atg	cat	ttg	cca	agt	cct	2976
294	Arg	Tyr	Leu	Val	Ile	Gln	Gly	Asp	Glu	Arg	Met	His	Leu	Pro	Ser	Pro	
295		980			985			990									
297	aca	gac	tcc	aac	tcc	tac	cgt	gcc	ctg	atg	gat	gaa	gaa	gac	atg	gac	3024
298	Thr	Asp	Ser	Asn	Phe	Tyr	Arg	Ala	Leu	Met	Asp	Glu	Glu	Asp	Met	Asp	
299		995			1000			1005									
301	gac	gtg	gtg	gat	gcc	gac	gag	tac	ctc	atc	cca	cag	cag	ggc	ttc		3069
302	Asp	Val	Val	Asp	Ala	Asp	Glu	Tyr	Leu	Ile	Pro	Gln	Gln	Gly	Phe		
303	1010			1015			1020										
305	ttc	agc	agc	ccc	tcc	acg	tca	cg	act	ccc	ctc	ctg	agc	tct	ctg		3114
306	Phe	Ser	Ser	Pro	Ser	Thr	Ser	Arg	Thr	Pro	Leu	Leu	Ser	Ser	Leu		
307	1025			1030			1035										
309	agt	gca	acc	agc	aac	aat	tcc	acc	gtg	gct	tgc	att	gat	aga	aat		3159
310	Ser	Ala	Thr	Ser	Asn	Asn	Ser	Thr	Val	Ala	Cys	Ile	Asp	Arg	Asn		
311	1040			1045			1050										
313	ggg	ctg	caa	agc	tgt	ccc	atc	aag	gaa	gac	agc	ttc	ttg	cag	cga		3204
314	Gly	Leu	Gln	Ser	Cys	Pro	Ile	Lys	Glu	Asp	Ser	Phe	Leu	Gln	Arg		
315	1055			1060			1065										
317	tac	agc	tca	gac	ccc	aca	ggc	gcc	ttg	act	gag	gac	agc	ata	gac		3249
318	Tyr	Ser	Ser	Asp	Pro	Thr	Gly	Ala	Leu	Thr	Glu	Asp	Ser	Ile	Asp		
319	1070			1075			1080										
321	gac	acc	ttc	ctc	cca	gtg	cct	gaa	tac	ata	aac	cag	tcc	gtt	ccc		3294
322	Asp	Thr	Phe	Leu	Pro	Val	Pro	Glu	Tyr	Ile	Asn	Gln	Ser	Val	Pro		
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327	1100			1105			1110										
329	cct	ctg	aac	ccc	gcg	ccc	agc	aga	gac	cca	cac	tac	cag	gac	ccc		3384
330	Pro	Leu	Asn	Pro	Ala	Pro	Ser	Arg	Asp	Pro	His	Tyr	Gln	Asp	Pro		
331	1115			1120			1125										
333	cac	agc	act	gca	gtg	ggc	aac	ccc	gag	tat	ctc	aac	act	gtc	cag		3429
334	His	Ser	Thr	Ala	Val	Gly	Asn	Pro	Glu	Tyr	Leu	Asn	Thr	Val	Gln		
335	1130			1135			1140										
337	ccc	acc	tgt	gtc	aac	agc	aca	ttc	gac	agc	cct	gcc	cac	tgg	gcc		3474
338	Pro	Thr	Cys	Val	Asn	Ser	Thr	Phe	Asp	Ser	Pro	Ala	His	Trp	Ala		
339	1145			1150			1155										
341	cag	aaa	ggc	agc	cac	caa	att	agc	ctg	gac	aac	cct	gac	tac	cag		3519
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VERIFICATION SUMMARY
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L:17 M:271 C: Current Filing Date differs, Replaced Current Filing Date
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